

## *Ins1* Gene Up-Regulated in a $\beta$ -Cell Line Derived from *Ins2* Knockout Mice

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The authors have derived a new  $\beta$ -cell line ( $\beta$ Ins2<sup>-/-lacZ</sup>) from *Ins2*<sup>-/-</sup> mice that carry the *lacZ* reporter gene under control of the *Ins2* promoter.  $\beta$ Ins2<sup>-/-lacZ</sup> cells stained positively using anti-insulin antibody, expressed  $\beta$ -cell-specific genes encoding the transcription factor PDX-1, glucokinase, and Glut-2, retained glucose-responsiveness for insulin secretion, and expressed the *lacZ* gene. Analysis of *Ins1* expression by reverse transcriptase–polymerase chain reaction (RT-PCR) showed that *Ins1* transcripts were significantly raised to compensate for the lack of *Ins2* transcripts in  $\beta$ Ins2<sup>-/-lacZ</sup> cells, as compared to those found in  $\beta$ TC1 cells expressing both *Ins1/Ins2*. Thus, transcriptional up-regulation of the remaining functional insulin gene in *Ins2*<sup>-/-</sup> mice could potentially contribute to the  $\beta$ -cell adaptation exhibited by these mutants, in addition to the increase in  $\beta$ -cell mass that we previously reported. We have also shown that *lacZ* expression, as analyzed by determining  $\beta$ -galactosidase activity, was up-regulated by incubating  $\beta$ Ins2<sup>-/-lacZ</sup> cells with GLP-1 and/or IBMX, 2 known stimulators of insulin gene expression. These cells thus represent a new tool for testing of molecules capable of stimulating *Ins2* promoter activity.

**Keywords**  $\beta$ -Cell Line; Compensatory Responses; Diabetes; Glucose Homeostasis; Insulin Genes; Knockout Mice

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Insulin resistance and  $\beta$ -cell dysfunction are the two major characteristic features of type 2 diabetes. It is now well admitted that overt diabetes does not develop as long as the  $\beta$ -cells can secrete increasing amounts of insulin to overcome insulin resistance [1]. Several groups have extensively investigated the molecular mechanisms that regulate insulin gene expression as well as insulin secretion during the past several years [2, 3]. Besides, a few studies in rodents also showed that the  $\beta$ -cell mass could significantly increase during pregnancy [4] or after glucose infusion for a short period, revealing potential plasticity of the  $\beta$ -cell compartment [5, 6]. Moreover, the study of knockout mice for insulin receptor substrate (IRS)-1, an intracellular mediator of insulin signaling, provided convincing evidence that insulin resistance could be overcome if  $\beta$ -cell mass were increased [7]. This conclusion was strongly supported by the observation that insulin resistance in IRS-2–knockout mice, which fail to increase their  $\beta$ -cell mass, leads to overt diabetes [8].

In recent years, transgenic and gene-targeting approaches in the mouse were extensively applied to examine the effects of manipulating the expression of genes encoding key players in  $\beta$ -cell development and/or function, such as the transcription factor PDX-1 [9, 10] or proteins involved in glucose-regulated insulin secretion, such as glucokinase (GK) [11] or the glucose transporter Glut-2 [12]. We previously generated knockout mice for the *Ins1* and *Ins2* genes [13]. The single *Ins1*<sup>-/-</sup> or *Ins2*<sup>-/-</sup> mutants were viable, fertile, and did not show any major metabolic alteration. Total pancreatic insulin content in *Ins1*<sup>-/-</sup> or *Ins2*<sup>-/-</sup> mice was comparable to that found in control animals, indicating that compensatory up-regulations take place in these mutants. Analysis of the

transcripts of the remaining functional insulin gene revealed a dramatic increase of *Ins1* transcripts in total pancreatic RNA from *Ins2*<sup>-/-</sup> mice. Interestingly, morphometric analysis of the pancreases showed that  $\beta$ -cell mass in both kinds of mutants was significantly augmented, particularly in *Ins2*<sup>-/-</sup> mutants. Thus, absence of either one of the 2 insulin genes was partly compensated at the tissue level by a specific increase of the  $\beta$ -cell compartment [13].

It was, however, difficult to determine whether some compensatory up-regulation of *Ins1* could also take place in individual  $\beta$ -cells in *Ins2*<sup>-/-</sup> mice. We have addressed this issue here using a  $\beta$ -cell line derived from *Ins2*<sup>-/-</sup> mutants. Because *Ins2*<sup>-/-</sup> mice carry *lacZ* reporter gene under the control of the *Ins2* promoter, such  $\beta$ -cells also represent an interesting tool for the testing of drugs that could up-regulate *Ins2* promoter activity.

## MATERIALS AND METHODS

### Animals

Generation of *Ins2*<sup>-/-</sup> mice has been described previously [14]. These mutant mice were crossed with Rip1-Tag2 mice [15] (kindly provided by D. Hanahan) and an *Ins2*<sup>-/-</sup>, Rip1-Tag2 mouse line was established. The offspring were genotyped by polymerase chain reaction (PCR) using specific primers.

### Derivation of the $\beta$ *Ins2*<sup>-/-lacZ</sup> Cell Line

$\beta$ -Cell lines from *Ins2*<sup>-/-</sup>, Rip1-Tag2 mice were isolated as previously described [16]. Briefly, 10-week-old *Ins2*<sup>-/-</sup>, Rip1-Tag2 mice were killed by cervical dislocation, and the pancreases were perfused through the bile duct with 2 mg/mL collagenase (Sigma, St. Louis, MO) dissolved in Hank's buffered saline solution (HBSS; Life Technologies, Gaithersburg, MD). The pancreases were incubated for 20 minutes at 37°C with gentle agitation. The digested material was washed 3 times in HBSS containing 10% fetal bovine serum (FBS; Techgen, Les Ullis, France) and then recovered in RPMI with 10% FBS. Islets were hand-picked under a dark-field microscope and cultured in the same medium in an incubator with 5% CO<sub>2</sub> at 37°C. Islets were then placed individually in 24-well plates and cultured in high-glucose medium containing 25 mM Hepes (Life Technologies), 15% horse serum (Life Technologies), 2.5% FBS, 200 U/mL penicillin, and 200  $\mu$ g/mL streptomycin. After approximately 2 months in culture, islets that clearly expanded were removed, trypsinized, and plated onto 96-well plates to recover  $\beta$ -cell lines. At confluency, cells were usually split at a 1:2 or 1:3 ratio.  $\beta$ TC1 cells (obtained from D. Hanahan) used in some experiments were cultured in the same medium.

## Immunocytochemistry and Histochemistry

For insulin detection, cells were fixed in 4% paraformaldehyde, incubated first with guinea pig polyclonal anti-insulin antibody and then with peroxidase-coupled goat anti-guinea pig serum as described [14]. The peroxidase activity was revealed using diaminobenzidine. To detect  $\beta$ -galactosidase ( $\beta$ -gal) activity, cells were fixed in 0.25% glutaraldehyde and then stained using 5-bromo-4-chloro-3-indolyl  $\beta$ -D-galactoside (X-gal) as described [13].

## Reverse Transcriptase (RT)-PCR Analysis of Gene Expression

The primers and probes used to analyze insulin 1/insulin 2, glucagon, somatostatin, pancreatic polypeptide (PP), and  $\beta$ -actin transcripts have been specified previously [13]. The primer sets and oligonucleotide probes used to analyze the transcripts for PDX-1, GK, Glut-2, and TATA-binding protein (TBP) were as follows: 5'-TCGCTGGGATCACTGGAGCA-3'/5'-GGTCCGCTGTGT AAGCACC-3' and 5'-[<sup>32</sup>P]-GACC TTTCCCGAATGGAA CC-3' for PDX-1; 5'-CACCCAAC TCGAAATCACC-3'/5'-CATTTGTGGGGAGTC-5' and 5'-[<sup>32</sup>P]-GGGCCAGTCAAATCCAGGCA-3' for GK; 5'-GA GCCAAGGACCCCGTCCTA-3'/5'-GTGAAGACCAGGAC CACCCC-3' and 5'-[<sup>32</sup>P]-GCCCTCTGCTTCCAGTACAT-3' for Glut-2; 5'-AAGAGAGCCACGGACAACCTG-3'/5'-TACT GAACTGCTGGTGGGTC-3' and 5'-[<sup>32</sup>P]-GAGTTGTGCA GAAGTTGGGC-3' for TBP. The following PCR reactions were used: 48°C, 45 minutes; 94°C, 2 minutes, 94°C, 30 seconds; 60°C, 1 minute, 68°C, 2 minutes for 25 cycles; 72°C 10 minutes. The PCR products were run on agarose gels, transferred onto Hybond membranes (Amersham, Saclay, France), and hybridized using [<sup>32</sup>P]-labeled oligonucleotide probes. For insulin 1/insulin 2, a unique primer pair was used for RT-PCR, and the PCR products were digested with MspI before Southern blot analysis. Hybridization using a single [<sup>32</sup>P]-labeled oligonucleotide probe revealed a fragment of 71 bp for insulin 1 and another of 112 bp for insulin 2. Quantification of PCR products was performed using a Phosphorimager equipped with ImageQuant software (Molecular Dynamics) and was expressed in relation to the internal control.

## Total Cellular Insulin Content and Insulin Secretion

Total cellular insulin content was obtained by radioimmunoassay (RIA) kit (ICN, Irvine, CA) using acid/alcohol extracts as described [16]. Insulin secretion assays were performed with cells cultured in 48-well plates when they reached 70% to 80% confluency as described [15]. Briefly, cells were preincubated for 1 hour in Krebs-Ringer buffer, then incubated

for 2 hours in Krebs-Ringer buffer containing different glucose concentrations as indicated. The secretion medium was recovered, centrifuged for 10 minutes at 1000 rpm, and supernatants were stored at  $-80^{\circ}\text{C}$  until used for determining insulin levels by RIA.

### Determination of $\beta$ -Gal Activity

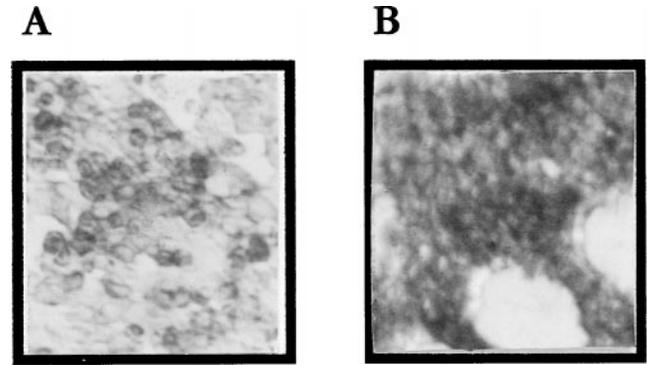
Cells were cultured in 48-well plates in high-glucose medium and incubated for 48 hours in the presence of GLP-1 (Sigma) and/or 3-isobutyl-1-methylxanthine (IBMX) (Sigma) at the indicated concentrations. Cells were then washed with phosphate-buffered saline (PBS) solution and lysed in a commercial lysis buffer (Boehringer, Mannheim, Germany).  $\beta$ -Gal activity in the protein extracts was determined by colorimetric assay using o-nitrophenyl  $\beta$ -D-galactopyranoside (ONPG) (Sigma) as substrate. Blank values obtained with lysates from  $\beta$ TTC1 cells were subtracted.

## RESULTS AND DISCUSSION

### $\beta$ -Cell Line Derived From $Ins2^{-/-}$ Mice

In previous studies,  $\beta$ -cell lines were isolated from islets of transgenic mice expressing SV40 large T antigen gene under the control of rat *Ins1* gene promoter (Rip1-Tag2 mice) [15, 16]. Therefore  $Ins2^{-/-}$  mice were crossed with Rip1-Tag2 mice and an  $Ins2^{-/-}$ , Rip1-Tag2 mouse line was established. Hyperplastic islets from pancreases of  $Ins2^{-/-}$ , Rip1-Tag2 mice were isolated prior to tumor formation, i.e., at 8 to 10 weeks. After a 2-month culture in high-glucose medium (22 mM), some islets clearly expanded as a result of efficient proliferation of  $\beta$ -cells transformed by SV40 T antigen, and individually trypsinized. One of the  $\beta$ -cell lines recovered from cultured cells used in this study was designated  $\beta Ins2^{-/-lacZ}$ . These cells showed a doubling time of 10 days and grew in islets-like clusters. The cells were not further subcloned to avoid clonal effects and used up to passage 19 in the experiments described here.

We first confirmed that  $\beta Ins2^{-/-lacZ}$  cells were  $\beta$ -cells.  $\beta Ins2^{-/-lacZ}$  cells were stained using an anti-insulin antibody (Figure 1A).  $Ins2^{-/-}$  mice carry the *lacZ* reporter gene under control of the *Ins2* promoter and the presence of  $\beta$ -gal in  $\beta Ins2^{-/-lacZ}$  cells was detected by X-gal staining (Figure 1B). Because the explanted islets of origin also contained  $\alpha$ ,  $\delta$ , and PP cells, which produce glucagon, somatostatin, and pancreatic polypeptide, respectively, we checked for the expression of the corresponding genes for these hormones by RT-PCR, using total RNA from  $\beta Ins2^{-/-lacZ}$  cells. No transcript was found for somatostatin and pancreatic polypeptide. Few glucagon transcripts were detected (not shown), as is fre-



**FIGURE 1**

Immunocytochemical detection of insulin (A) and histochemical analysis of *lacZ* expression (B) in  $\beta Ins2^{-/-lacZ}$  cells. (A) Cells were fixed and stained using an anti-insulin antibody. (B) Cells were fixed and  $\beta$ -gal activity was visualized by X-gal staining. Magnification:  $10\times$ .

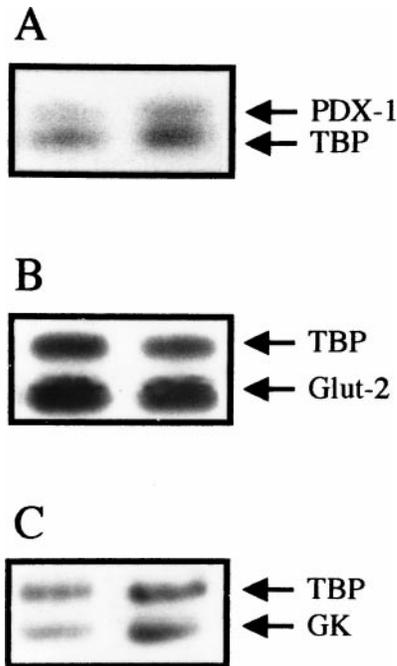
quently observed when deriving a  $\beta$ -cell line using Rip1-Tag2 construct [17].

We also analyzed the expression of genes encoding PDX-1, GK, and Glut-2. PDX-1 is a transcription factor that is essential for pancreas development and in the regulation of islet gene expression in mature  $\beta$ -cells [18]. It was shown that PDX-1 expression in the liver upon adenoviral vector-mediated gene transfer in the mouse resulted in transdifferentiation of a small fraction of hepatocytes into insulin-secreting cells [19]. GK is the high- $K_M$  hexokinase present in both  $\beta$ -cells as well as in hepatocytes and catalyzes the initial conversion of glucose to glucose-6-phosphate. Finally, Glut-2 is a glucose transporter also present in  $\beta$ -cells and hepatocytes [20]. The role of GK and Glut-2 in glucose sensing was fully confirmed by global and tissue-specific disruption of the corresponding genes in the mouse. Knockout mice for GK or Glut-2 presented defective insulin secretion and developed diabetes [11, 12]. As presented in Figure 2, transcripts for PDX-1, GK, and Glut-2 could be readily detected by RT-PCR using total RNA from  $\beta Ins2^{-/-lacZ}$  cells.

In conclusion,  $\beta Ins2^{-/-lacZ}$  cells continue to synthesize insulin and express important  $\beta$ -cell-specific marker genes. Moreover, *Ins2* promoter drives expression of *lacZ* gene in these cells.

### Glucose-Induced Insulin Secretion From $\beta Ins2^{-/-lacZ}$ Cells

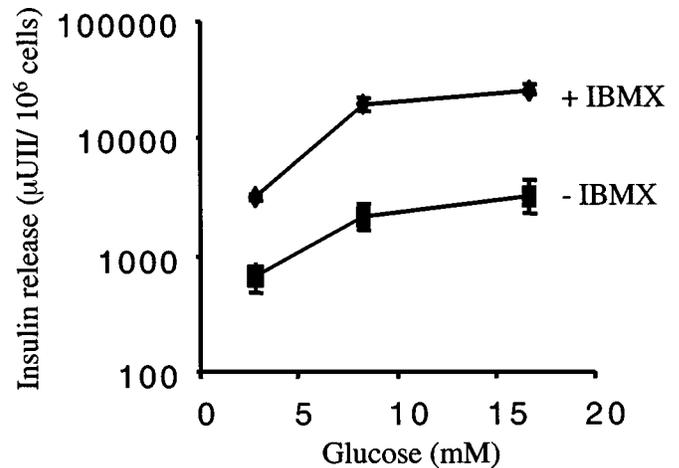
We examined the ability of  $\beta Ins2^{-/-lacZ}$  cells to secrete insulin in response to glucose because this property is frequently lost when deriving a  $\beta$ -cell line [17]. We first measured total cellular insulin content in acid-alcohol extracts



**FIGURE 2**

RT-PCR analysis of  $\beta$ -cell-specific gene expression. Transcripts for PDX-1 (A), Glut-2 (B), and GK (C) were amplified using total RNA from  $\beta$ Ins2<sup>-/-lacZ</sup> cells and analyzed by Southern blotting. TBP mRNA was coamplified as control. The results of 2 separate samples are presented. The sizes of various PCR products correspond to 275 bp (PDX-1), 150 bp (Glut-2), 162 bp (GK), and 233 bp (TBP).

of  $\beta$ Ins2<sup>-/-lacZ</sup> cells by RIA. The values obtained ( $30.04 \pm 2.167$  mUI/10<sup>6</sup> cells) were comparable to those previously reported with other murine  $\beta$ -cell lines in which both the *Ins1* and *Ins2* genes were functional [15, 16]. Thus, increased *Ins1* gene expression appears to quantitatively compensate the absence of insulin transcripts from *Ins2*-null alleles in  $\beta$ Ins2<sup>-/-lacZ</sup> cells. Glucose-induced insulin secretion from  $\beta$ Ins2<sup>-/-lacZ</sup> cells was subsequently analyzed by incubating the cells in Krebs-Ringer secretion buffer supplemented with 2.8, 8.3, or 16.7 mM glucose. After incubation for 2 hours, the amount of insulin released in the medium was determined by RIA and the results obtained are presented in Figure 3. Insulin release from  $\beta$ Ins2<sup>-/-lacZ</sup> cells increased with increasing glucose concentrations. The values obtained at 16.7 mM glucose were about 5-fold higher as compared to those at 2.8 mM glucose. Insulin secretion from  $\beta$ Ins2<sup>-/-lacZ</sup> cells was also examined in the presence of IBMX, which is known to potentiate glucose-stimulated insulin release from  $\beta$ -cells. Addition of 0.5 mM IBMX in the secretion buffer enhanced insulin release from  $\beta$ Ins2<sup>-/-lacZ</sup> cells at all glucose concen-



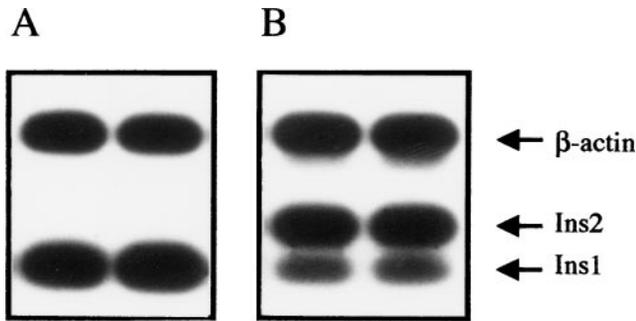
**FIGURE 3**

Glucose-induced insulin secretion in  $\beta$ Ins2<sup>-/-lacZ</sup> cells. Insulin release was measured in the absence or in the presence of 0.5 mM IBMX. The incubation time was 2 hours. Values are means  $\pm$  SE from triplicate assays.

trations (Figure 3). In the presence of IBMX, insulin secretion at high glucose concentration (16.7 mM) was  $\sim$ 8.3-fold higher than that at low glucose concentration (2.8 mM). All these results indicate that  $\beta$ Ins2<sup>-/-lacZ</sup> cells have retained glucose responsiveness for insulin secretion.

### *Ins1* Gene is Up-Regulated in $\beta$ Ins2<sup>-/-lacZ</sup> Cells

To investigate possible up-regulation of the *Ins1* gene in the absence of *Ins2* transcripts in  $\beta$ Ins2<sup>-/-lacZ</sup> cells, we compared insulin transcripts produced in  $\beta$ Ins2<sup>-/-lacZ</sup> cells with those in  $\beta$ TC1 cells. This latter cell line was derived from Rip1-Tag2 mice in which both the *Ins1* and *Ins2* genes are functional. Insulin gene expression was analyzed by RT-PCR using total cellular RNA. In  $\beta$ TC1 cells, *Ins1* transcripts represent the minority of insulin transcripts (Figure 4B). In  $\beta$ Ins2<sup>-/-lacZ</sup> cells, the amounts of *Ins1* transcripts were significantly higher than those found in  $\beta$ TC1 cells (Figure 4A). Quantitative analysis of RT-PCR products for insulin as well as  $\beta$ -actin transcripts showed that *Ins1* transcripts in  $\beta$ Ins2<sup>-/-lacZ</sup> cells represent 55% to 88% of total insulin transcripts present in  $\beta$ TC1 cells. These results indicate that the *Ins1* gene is up-regulated in  $\beta$ Ins2<sup>-/-lacZ</sup> cells to compensate for the absence of *Ins2* transcripts. It can be inferred from this study that *Ins1* gene up-regulation could also potentially take place in individual  $\beta$ -cells in vivo in *Ins2*<sup>-/-</sup> mice, in the compensating responses exhibited by these mutants to maintain normal glucose homeostasis [14].



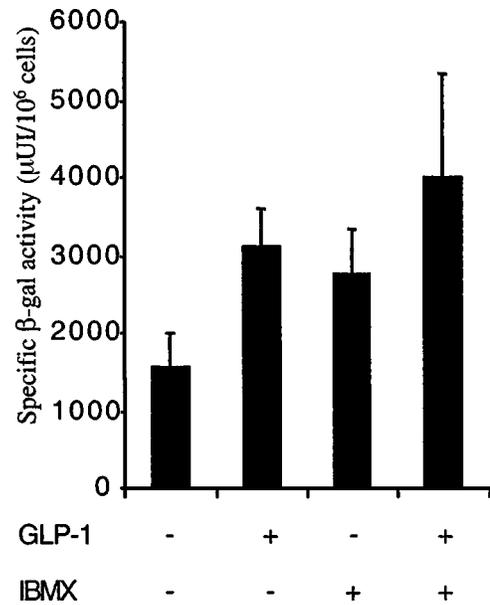
**FIGURE 4**

RT-PCR analysis of insulin gene expression in  $\beta$ Ins2<sup>-/-lacZ</sup> (A) and  $\beta$ TC1 (B) cells. Transcripts for Insulin 1 (Ins1)/Insulin 2 (Ins2) were amplified using total RNA from  $\beta$ Ins2<sup>-/-lacZ</sup> or  $\beta$ TC1 cells and analyzed by Southern blotting.  $\beta$ -Actin mRNA was coamplified as control. The results of 2 separate samples are presented. The sizes of various PCR products correspond to 76 bp (Insulin 1), 112 bp (Insulin 2), and 243 bp ( $\beta$ -actin).

### Induction of the *Ins2-lacZ* Gene in $\beta$ Ins2<sup>-/-lacZ</sup> Cells

Regulation of insulin gene expression has been examined in many studies using different  $\beta$ -cell lines transfected with reporter genes under control of rodent or human insulin promoter. In some of these studies, several-fold induction of insulin promoter activity by glucose was reported [21, 22]. Because  $\beta$ Ins2<sup>-/-lacZ</sup> cells express the *lacZ* gene inserted at the *Ins2* locus, these cells represent an interesting tool for testing or screening of molecules that can up-regulate *Ins2* promoter activity, because *lacZ* expression can be easily monitored by measuring  $\beta$ -gal activity in cellular extracts. We could not examine the effect of glucose on *Ins2-lacZ* expression, because only a moderate decrease, if any, of  $\beta$ -gal activity was observed when  $\beta$ Ins2<sup>-/-lacZ</sup> cells grown in high glucose were transferred to low-glucose medium (not shown). We could, however, test the effect of GLP-1 and IBMX on *Ins2-lacZ* gene expression in  $\beta$ Ins2<sup>-/-lacZ</sup> cells. GLP-1 is a hormone secreted by the gut during digestion and has been shown to stimulate insulin gene transcription under hyperglycemic conditions [23]. IBMX has also been reported to stimulate insulin gene expression [24].

As shown in Figure 5, incubation of  $\beta$ Ins2<sup>-/-lacZ</sup> cells cultured in high-glucose medium with 10<sup>-7</sup> M GLP-1 for 2 days resulted in 97% increase in  $\beta$ -gal activity as compared to  $\beta$ Ins2<sup>-/-lacZ</sup> cells cultured without GLP-1. A similar increase in  $\beta$ -gal activity was also observed for  $\beta$ Ins2<sup>-/-lacZ</sup> cells incubated with 0.5 mM IBMX. The  $\beta$ -gal activity in  $\beta$ Ins2<sup>-/-lacZ</sup> cells cultured in the presence of both GLP-1 and IBMX was increased up to 157% as compared to untreated



**FIGURE 5**

Analysis of  $\beta$ -gal activity in  $\beta$ Ins2<sup>-/-lacZ</sup> cells cultured under various conditions.  $\beta$ -Gal activity was determined in  $\beta$ Ins2<sup>-/-lacZ</sup> cells cultured in high-glucose medium (22 mM) supplemented or not for 48 hours with 10<sup>-7</sup> M GLP-1 and/or 0.5 mM IBMX. A blank value obtained with  $\beta$ TC1 cells was subtracted. Values are means  $\pm$  SE from triplicate and quadruplicate assays.

cells. These results demonstrate that the *Ins2* promoter can be induced in  $\beta$ Ins2<sup>-/-lacZ</sup> cells.

In conclusion, the advantage of such a model  $\beta$ Ins2<sup>-/-lacZ</sup> cell line is twofold: (1) to understand the transcriptional network governing the expression of the *Ins2* promoter, and (2) to study the mechanisms involved in the up-regulation of *Ins1* promoter activity. These cells also represent an interesting new tool for the screening of molecules that could stimulate *Ins2* promoter activity under high-glucose conditions, and therefore would have potential therapeutic interest for type 2 diabetes, because the *Ins2* gene in the mouse is homologous to the human insulin gene.

### REFERENCES

- [1] Kahn, B. B. (1998) Type 2 diabetes: When insulin secretion fails to compensate for insulin resistance. *Cell*, **92**, 593–596.
- [2] Docherty, K., and Clark, A. R. (1994) Nutrient regulation of insulin gene expression. *FASEB J.*, **8**, 20–27.
- [3] Dumonteil, E., and Philippe, J. (1996) Insulin gene: Organisation, expression and regulation. *Diabetes Metab.*, **22**, 164–173.
- [4] Nieuwenhuizen, A. G., Schuiling, G. A., Moes, H., and Koiter, T. R. (1997) Role of increased insulin demand in the adaptation of the endocrine pancreas to pregnancy. *Acta Physiol. Scand.*, **159**, 303–312.

- [5] Bernard, C., Thibault, C., Berthault, M. F., Magnan, C., Saulnier, C., Portha, B., Pralong, W. F., Penicaud, L., and Ktorza, A. (1998) Pancreatic beta-cell regeneration after 48-h glucose infusion in mildly diabetic rats is not correlated with functional improvement. *Diabetes*, **47**, 1058–1065.
- [6] Bonner, W. S., Deery, D., Leahy, J. L., and Weir, G. C. (1989) Compensatory growth of pancreatic beta-cells in adult rats after short-term glucose infusion. *Diabetes*, **38**, 49–53.
- [7] Araki, E., Lipes, M. A., Patti, M. E., Bruning, J. C., Haag, B. R., Johnson, R. S., and Kahn, C. R. (1994) Alternative pathway of insulin signalling in mice with targeted disruption of the IRS-1 gene. *Nature*, **372**, 186–190.
- [8] Withers, D. J., Gutierrez, J. S., Towery, H., Burks, D. J., Ren, J. M., Previs, S., Zhang, Y., Bernal, D., Pons, S., Shulman, G. I., Bonner, W. S., and White, M. F. (1998) Disruption of IRS-2 causes type 2 diabetes in mice. *Nature*, **391**, 900–904.
- [9] Jonsson, J., Carlsson, L., Edlund, T., and Edlund, H. (1994) Insulin-promoter-factor 1 is required for pancreas development in mice. *Nature*, **371**, 606–609.
- [10] Marshak, S., Ben-Shushan, E., Shoshkes, M., Havin, L., Cerasi, E., and Melloul, D. (2001) Regulatory elements involved in human pdx-1 gene expression. *Diabetes*, **50(Suppl 1)**, S37–S38.
- [11] Bali, D., Svetlanov, A., Lee, H. W., Fusco-DeMane, D., Leiser, M., Li, B., Barzilai, N., Surana, M., Hou, H., and Fleischer, N. (1995) Animal model for maturity-onset diabetes of the young generated by disruption of the mouse glucokinase gene. *J. Biol. Chem.*, **270**, 21464–21467.
- [12] Guillam, M. T., Hummler, E., Schaerer, E., Yeh, J. I., Birnbaum, M. J., Beermann, F., Schmidt, A., Deriaz, N., Thorens, B., and Wu, J. Y. (1997) Early diabetes and abnormal postnatal pancreatic islet development in mice lacking Glut-2. *Nat. Genet.*, **17**, 327–330.
- [13] Leroux, L., Desbois, P., Lamotte, L., Duvillié, B., Cordonnier, N., Jackerott, M., Jami, J., Bucchini, D., and Joshi, R. L. (2001) Compensatory responses in mice carrying a null mutation for Ins1 or Ins2. *Diabetes*, **50(Suppl 1)**, S150–S153.
- [14] Duvillié, B., Cordonnier, N., Deltour, L., Dandoy, Dron F., Itier, J. M., Monthieux, E., Jami, J., Joshi, R. L., and Bucchini, D. (1997) Phenotypic alterations in insulin-deficient mutant mice. *Proc. Natl. Acad. Sci. U.S.A.*, **94**, 5137–5140.
- [15] Efrat, S., Leiser, M., Surana, M., Tal, M., Fusco-DeMane, D., and Fleischer, N. (1993) Murine insulinoma cell line with normal glucose-regulated insulin secretion. *Diabetes*, **42**, 901–907.
- [16] Radvanyi, F., Christgau, S., Baekkeskov, S., Jolicoeur, C., and Hanahan, D. (1993) Pancreatic beta cells cultured from individual preneoplastic foci in a multistage tumorigenesis pathway: A potentially general technique for isolating physiologically representative cell lines. *Mol. Cell. Biol.*, **13**, 4223–4232.
- [17] Levine, F. (1997) Gene therapy for diabetes: Strategies for beta-cell modification and replacement. *Diabetes Metab. Rev.*, **13**, 209–246.
- [18] Yamaoka, T., and Itakura, M. (1999) Development of pancreatic islets. *Int. J. Mol. Med.*, **3**, 247–261.
- [19] Ferber, S., Halkin, A., Cohen, H., Ber, I., Einav, Y., Goldberg, I., Barshack, I., Seiffers, R., Kopolovic, J., Kaiser, N., and Karasik, A. (2000) Pancreatic and duodenal homeobox gene 1 induces expression of insulin genes in liver and ameliorates streptozotocin-induced hyperglycemia. *Nat. Med.*, **6**, 568–572.
- [20] Rolland, F., Winderickx, J., and Thevelein, J. M. (2001) Glucose-sensing mechanisms in eukaryotic cells. *Trends Biochem. Sci.*, **26**, 310–317.
- [21] de Vargas, L., Sobolewski, J., Siegel, R., and Moss, L. G. (1997) Individual beta cells within the intact islet differentially respond to glucose. *J. Biol. Chem.*, **272**, 26573–26577.
- [22] Leibiger, B., Moede, T., Schwarz, T., Brown, G. R., Kohler, M., Leibiger, I. B., and Berggren, P. O. (1998) Short-term regulation of insulin gene transcription by glucose. *Proc. Natl. Acad. Sci. U.S.A.*, **95**, 9307–9312.
- [23] Drucker, D. J. (1998) Glucagon-like peptides. *Diabetes*, **47**, 159–169.
- [24] Kohnert, K. D., Ziegler, B., Hahn von Dorsche, H., Hehmke, B., and Schroder, D. (1982) Effects of 3-isobutyl-1-methylxanthine on neonatal pancreatic islets maintained in tissue culture. *Mol. Cell. Endocrinol.*, **28**, 425–437.